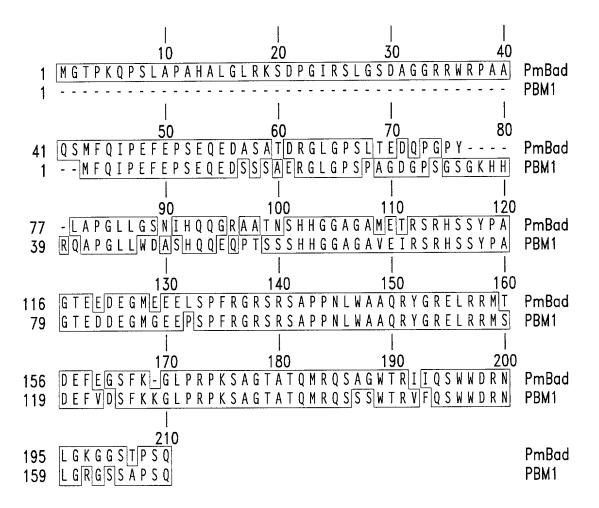
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Fig. 1A

AGC TCC GCC CCC TCC CAG TGACCTTCGG TCCACATCCC GAAATCCACC 585 Ser Ser Ala Pro Ser Gln 165 645 CGTTCCCATT GCCCTGGGCA GCCATTTTGA ATATGGGAGG AAGTAAGTTC CCTCAGGCCT 705 ATGCAAAAAG AGGATCCGTG CTGTATCCTT TGGAGGGAGG GTTGACCCAG ATTCCCTTCC GGTGTGTGT AAGCCACGGA AGGTTGGTCC CATCGGAAGT TTTGGGTTTT CCGCCCACAG 765 CCGCCGGAAG TGGCTCCGTG GCCCCGCCCT CAGGTTCCGG GGTTTCCCCC AGGCGCCTGC 825 GCTAAGTAGC GAGCCAGGTT TAACCGTTGT GTCACCGGGA CCCGAGCCCC CGCGATGCCC 885 TGGGGGCCGT GATCAGTACC AAATGTTAAT AAAGCCCGCG TGTGTGCCAA AAAAAAAAA 945 946

Fig. 1B

ALIGNMENT OF MOUSE AND HUMAN BAD PROTEIN SEQUEUNCES



PmBad = mouse protein sequence PBM1 = human protein sequence

Fig. 2

Bad 2-Hybrid Interactions

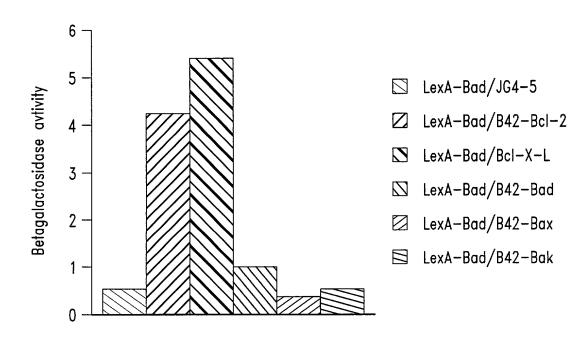


Fig. 3

Inventor(s): William A. Horne et al.

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Docket No. 480140.428D3

Binding of Bcl-2 Family Proteins to Bad

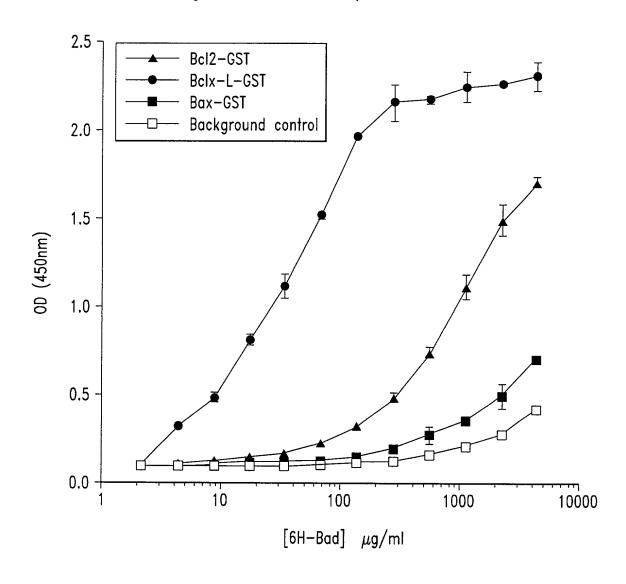


Fig. 4